

Supplementary Fig. 3: Validation of differential gene expression by quantitative real-time PCR. (a-b) To validate differential expression in NKG2DLneg versus NKG2DLpos subpopulations, qRT-PCR analysis was performed on genes found up- (a) or down-regulated (b) in the microarray data set; shown are results of n=3 AML (#8, #6, #1), analysed each in technical triplicates; the dotted line represents gene expression in NKG2DLpos subpoulations. (c) qRT-PCR verification of the 17 gene stemness score; analysis was performed in the same manner as in a-b.

